P. Gambel

PAGE: 1

RAW SEQUENCE LISTING PATENT APPLICATION US/08/753,851

1754

DATE: 02/03/2000

TIME: 11:13:49

INPUT SET: S34624.raw

This Raw Listing contains the General Information Section and up to the first 5 pages.

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1
                                       SEQUENCE LISTING
 2
                                                         ENTERED
            General Information:
 3
     (1)
          (i) APPLICANT: WEINBERG, J. BRICE
 5
                         HAYNES, BARTON F.
 7
         (ii) TITLE OF INVENTION: AN ADHESION MOLECULE
 9
        (iii) NUMBER OF SEQUENCES: 16
10
1.1
         (iv) CORRESPONDENCE ADDRESS:
12
13
               (A) ADDRESSEE: NIXON & VANDERHYE P.C.
14
               (B) STREET: 1100 NORTH GLEBE ROAD, 8TH FLOOR
15
               (C) CITY: ARLINGTON
               (D) STATE: VIRGINIA
16
17
               (E) COUNTRY: U.S.A.
18
               (F) ZIP: 22201-4714
19
20
          (v) COMPUTER READABLE FORM:
               (A) MEDIUM TYPE: Floppy disk
21
22
               (B) COMPUTER: IBM PC compatible
23
               (C) OPERATING SYSTEM: PC-DOS/MS-DOS
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
25
         (vi) CURRENT APPLICATION DATA:
26
27
               (A) APPLICATION NUMBER: 08/753,851
               (B) FILING DATE:
28
               (C) CLASSIFICATION:
29
30
31
       (vii) PRIOR APPLICATION DATA:
32
               (A) APPLICATION NUMBER: US 08/047,068
33
               (B) FILING DATE: 16-APR-1993
34
35
               (A) APPLICATION NUMBER: US 07/945,581
               (B) FILING DATE: 16-SEP-1992
36
37
       (vii) PRIOR APPLICATION DATA:
38
39
               (A) APPLICATION NUMBER: US 07/682,518
40
               (B) FILING DATE: 09-APR-1991
41
42
       (vii) PRIOR APPLICATION DATA:
43
               (A) APPLICATION NUMBER: US 07/669,730
44
               (B) FILING DATE: 15-MAR-1991
45
      (viii) ATTORNEY/AGENT INFORMATION:
46
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# RAW SEQUENCE LISTING PATENT APPLICATION US/08/753,851

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51 52 53 54 55		(ix)	(A) (B)	TEI TEI	EPHC EFAX	NE: (7	N INE (703) 03) 8 97 N	81 316-	6-40 4100	00							
57 58	(2)	INFO	RMATI	ON F	FOR S	SEQ I	D NO:	:1:									
59 60 61 62 63		(i)	(A) (B)	LEN TYP	IGTH: PE: a	22 mino	ERIST amino ació inear	ac i									
64 65 66		(ii)	MOLE	CULE	TYF	E: p	eptio	de									
68 69		(xi)	SEQU	JENCE	DES	CRIP	TION:	: SE	Q II	ОИО	:1:						
70 71 72		Cys 1	Glu	Lys	Asn	Gly 5	Arg 1	ſyr	Ser	Ile	Ser 10	Arg	Thr	Glu	Ala	Ala 15	Asp
73 74 75		Cys	Cys	Lys	Ala 20	Phe	Asn <sub>.</sub>										
76 77	(2)	INFO	RMATI	ON F	OR S	EQ I	D NO:	2:									
78 79 80 81 82		(i)	(A) (B)	LEN TYP	IGTH: PE: a	23 mino	ERIST amino acid inear	ac i									
83 84 85 86		(ii)	MOLE	CULE	TYP	E: p	eptid	le									
87 88		(xi)	SEQU	ENCE	DES	CRIP	TION:	SE	Q II	NO:	:2:						
89 90 91		Cys 1	Asn	Thr	Ser	Gln 5	Tyr A	Asp	Thr	Tyr	Cys 10	Phe	Asn	Ala	Ser	Ala 15	Pro
92 93 94		Pro	Glu		Asp 20	Cys	Thr S	Ser									
95 96	(2)	INFOR	RMATI	ON F	OR S	EQ I	D NO:	3:									
97 98 99		(i)	(A)	LEN	GTH:	32	ERIST aminc acid	ac									

# RAW SEQUENCE LISTING PATENT APPLICATION US/08/753,851

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			•	INPUT SET: S3462	24.1
100			(D) TOPOLOGY: linear		
101			<u></u>		
102		(ii)	MOLECULE TYPE: peptide		
103					
104					
105		121	CROWDIGE BECCRETON CHO TO NO 3		
106		(X1)	SEQUENCE DESCRIPTION: SEQ ID NO:3:		
107		C++0	Arg Asp Gly Thr Arg Tyr Val Gln Lys Gly Glu	Tur Ara Thr Ass	_
108 109		Cys 1	5 10	15	.1
110		_	3	13	
111		Pro	Glu Asp Ile Tyr Pro Ser Asn Pro Thr Asp Asp	Asp Val Ser Sei	r
112		110	20 25	30	
113					
114					
115	(2)	INFO	RMATION FOR SEQ ID NO:4:		
116					
117		(i)	SEQUENCE CHARACTERISTICS:		
118			(A) LENGTH: 42 amino acids		
119			(B) TYPE: amino acid		
120			(D) TOPOLOGY: linear		
121					
122		(11)	MOLECULE TYPE: peptide		
123					
124					
125 126		(vi)	SEQUENCE DESCRIPTION: SEQ ID NO:4:		
127		(XI)	SEQUENCE DESCRIPTION: SEQ ID NO.4.		
128		Cvs	Arg Asp Gly Thr Arg Tyr Val Gln Lys Gly Glu	Tvr Arg Ile Ası	n
129		1	5 10	15	-
130		_			
131		Pro	Glu Asp Ile Tyr Pro Ser Asn Pro Thr Asp Asp	Asp Val Ser Ser	r
132			20 25	30	
133					
134		Gly	Ser Ser Ser Glu Arg Ser Ser Thr Ser		
135			35 40	•	
136					
137	(2)	INFO	RMATION FOR SEQ ID NO:5:		
138		( = )	GEOVERNOE CHARACMEDICATION		
139		(1)	SEQUENCE CHARACTERISTICS:		
140 141			<ul><li>(A) LENGTH: 21 amino acids</li><li>(B) TYPE: amino acid</li></ul>		
142			(D) TOPOLOGY: linear		
143			(b) forobodi. Timear		
144		(ii)	MOLECULE TYPE: peptide		
145		,	E CE COLOR DE CENTRA		
146					
147					
148		(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:5:		
149					
150		Cys	Tyr Arg Thr Asn Pro Glu Asp Ile Tyr Pro Ser		Ç
151		1	5 10	15	
152					

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Asp Asp Val Ser Ser
153
154
                       20
155
     (2) INFORMATION FOR SEQ ID NO:6:
156
157
          (i) SEQUENCE CHARACTERISTICS:
158
159
                (A) LENGTH: 21 amino acids
                (B) TYPE: amino acid
160
                (D) TOPOLOGY: linear
161
162
          (ii) MOLECULE TYPE: peptide
163
164
165
166
167
          (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:
168
169
          Cys Thr Val His Pro Ile Pro Asp Glu Asp Ser Pro Trp Ile Thr Asp
170
                           5
171
172
          Ser Thr Pro Arg Ile
173
                       20
174
175
    (2) INFORMATION FOR SEQ ID NO:7:
176
177
          (i) SEQUENCE CHARACTERISTICS:
178
               (A) LENGTH: 21 amino acids
179
                (B) TYPE: amino acid
               (D) TOPOLOGY: linear
180
181
          (ii) MOLECULE TYPE: peptide
182
183
184
185
186
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:
187
188
          Asp Ser Pro Trp Ile Thr Asp Ser Thr Asp Arg Ile Phe Ala Thr Arg
189
                           5
                                               10
190
          Asp Gln Asp Thr Ile
191
192
193
     (2) INFORMATION FOR SEQ ID NO:8:
194
195
196
          (i) SEQUENCE CHARACTERISTICS:
               (A) LENGTH: 35 amino acids
197
               (B) TYPE: amino acid
198
               (D) TOPOLOGY: linear
199
200
         (ii) MOLECULE TYPE: peptide
201
202
203
204
205
         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:
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207		_	Ala	Thr	Arg		GIn	Asp	Thr	Pne		Pro	ser	GLY	GLY		Hıs
208		1				5					10					15	
209		_	_				_					_	_		_	_	_
210		Thr	Thr	His	Glu	Ser	Glu	Asp	Gly	His	Ser	His	Gly	Ser		Glu	Gly
211					20					25					30		
212																	
213		Gly	Ala	Asn													
214				35													
215																	
216	(2)	INFO	RMAT:	ION 1	FOR S	SEQ :	ID NO	0:9:									
217																	
218		(i)	SEQU	JENCI	E CHA	ARAC	reris	STICS	3:								
219		(A) LENGTH: 13 amino acids															
220		(B) TYPE: amino acid															
221		(D) TOPOLOGY: linear															
222		(D) TOPOLOGI. IIMERI															
223		(44)	MOTE	20111 1	י ייעו	)E. *	oot i	. d.o.									
		(ii)	MOTTE	20111	5 111	- E: F	pept.	Lue									
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227		(xi)	SEQU	JENC	E DES	SCRII	OITS	1: SI	EQ II	ONO:	9:						
228				•	_	_											
229		Cys	Arg	Asp	Gly	Ile	Arg	Tyr	Val	Gln	Lys	Gly	Glu	Tyr			
230		1				5					10						
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232	(2)	INFO	RMATI	ON I	FOR S	SEQ ]	D NO	10:	:								
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234		(i)	SEQU	JENCI	E CHA	ARAC'	CERIS	STICS	3:								
235			(A)	LEN	GTH:	: 7 a	mino	aci	lds								
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243		(xi)	SEOU	ENCE	e DES	CRTE	TON	ı. SE	O TE	NO.	10.						
244		(22.1)	DIQU	LIVEL	. DEL	CICI	1101		1Q 11	, 110.	10.			•			
245		Dro	Ser	Λen	Dro	Thr	λen	Λen									
246			PET	ASII	FIU	5	Asp	Asp									
247		1				5											
247	(2)	TNEOT	7848 MT	-ONT T	7OD 0	100 7	-D NC										
	(2)	INFOF	(MAT.T	ON F	OR S	EQ 1	טוו ע.	): TT:									
249		123	GEOT			D 3 GG		.m 0.0									
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